FIG. 1

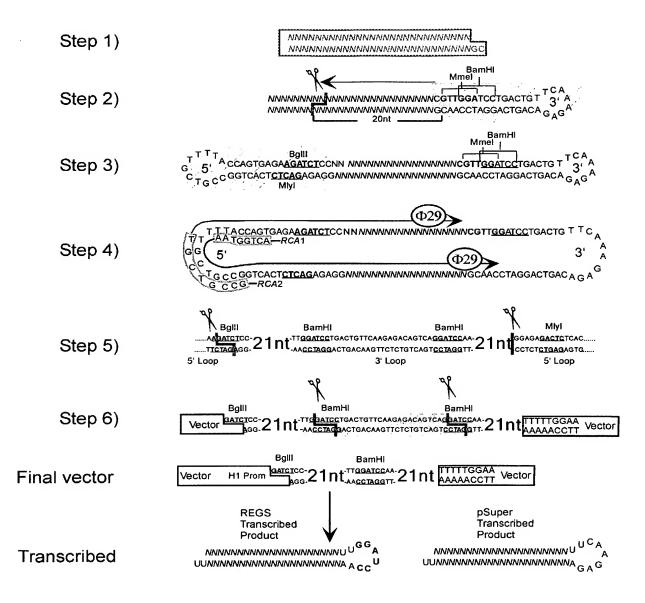
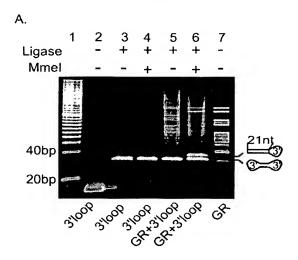
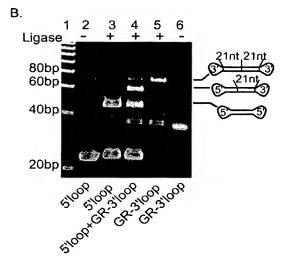


FIG. 2





C.

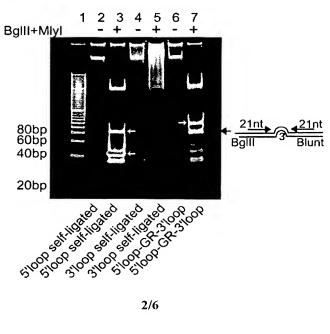


FIG. 3

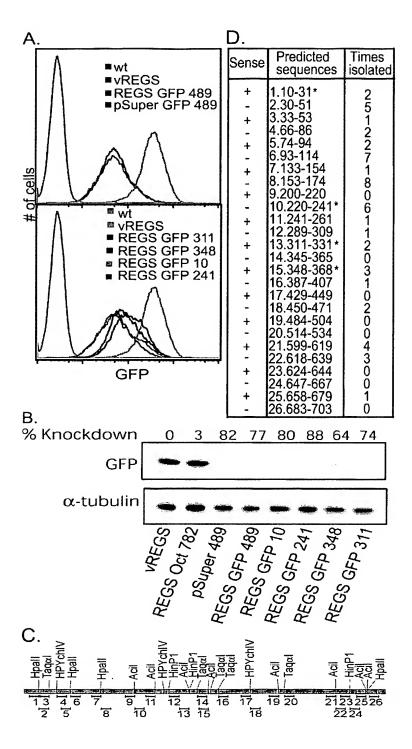


FIG. 4

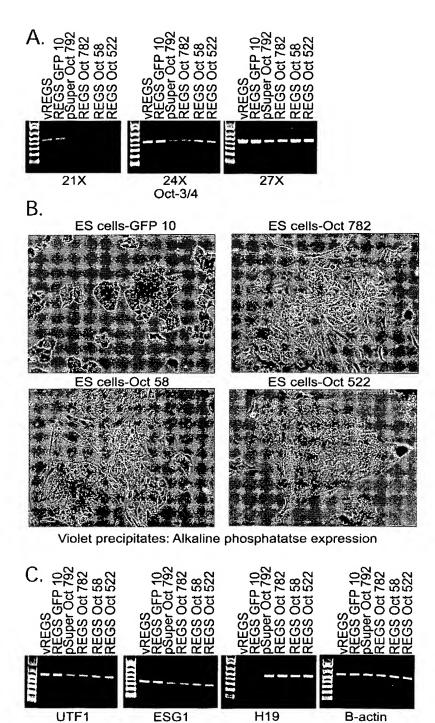
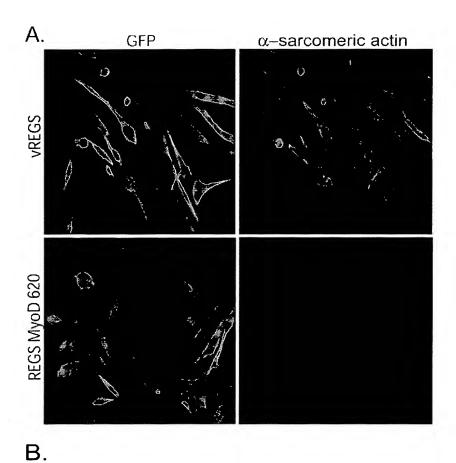


FIG. 5



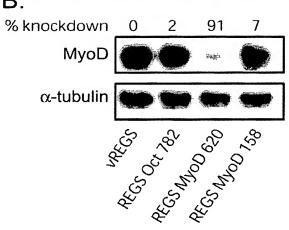


FIG. 6

siRNA location	Gene	Accession number
2966-2985 -	ATP7a	AK040753.1
278-258 -	Pleiotrophin	BC002064.1
3286-3269 -	RP23-263E3	AC122295.4
6296-6275 -	mitochondrial	V00711.1 MTCMM
493-473 -	GABA(A) receptor-associated protein 2	NM 026693.2
1465-1445 -	IMAGE:4527839 Ubiquitin conj enz homolog	BC016203.1
621-601 -	Ferritin heavy chain	B0012314.1
2427-2407 -	IMAGE:6489956	BC049362.1
32043225+	exportin 7	BC029702.1
805-785 -	Bat5	NM 178592.2
415-435+	EEF-2	BC002233.1
10533-10553+	CONTRACTOR THE GALLEGE CONTRACTOR OF THE PROPERTY OF A STATE OF THE PARTY OF THE PA	X14061.1 MMBGCXD
2142-2122 -	CDC46	D26090.1 MUSCDC46
	granded states and second and a second secon	BC040404.1
2761-2741 -	Angiotensin converting enzyme	BC046513.1
7484-7505 + 1549-1529 -	Fatty acid synthase IMAGE:5355806 EST	BC024847.1
450-471 +	CaMKII gamma	NM 178597.2
739-760 +	ldb3	NM 008321.1
1581-1601 +	PI3K subunit p85 alpha	BC051106.1
2771-2791 +	ESTIMAGE:3491837	AC123871.4
1147-1126 -	IMAGE:4194837	BC029813.1
	enolase 1	BC024644.1
1138-1158 +	F-box only protein	BC034854.1
78-58 -	similar to ribosomal L7a	XM 193935.1
951-972+	proprotein convertase subtilisin/kexin type 5	BC013068.1
758-779+	Krt2-5_	NM 027011.1
347-367+	Tubulin alpha 6	BC022182.1
265-244 -	Pouff1	NM 013633.1
488-509+	alpha-2-HS-glycoprotein	BC019822.1
1625-1605 -	prolyl 4 hydroxylase, beta polypeptide	BC008549.1
497-518 +	guanine nucleotide binding protein	BC048834.1
3588-3567 -	ubiquitin specific protease 7	BC046963.1
768-748 -	Diprp	NM 010088.1
2263-2284+	similar to pol protein	XM 196572.2
1437-1417 -	Sept3	AC104325.28
891-871 -	IMAGE:1515563	AK014200.1
2002-2022 +	RIKEN cDNA C230075L19	BC048924.1
1271-1252 -	glutathione peroxidase 3 (Gpx3)	NM 008161.1
2003-2024+	MLV-related provinus NL1 integrase	AY219562.2
13-33+	serum/glucocorticoid regulated kinase	BC005720.1
1044-1064 +	Ca2+-dependent ER nucleoside diphosphatase	BC020003.1
2824-2846+	Similar to cyclin Lania-6a	BC023747.1
	50 Total	
22 Antisense	3 contaminating plasmid sequence	especial for the first term.
20 Sense	3 no exact match	
	2 no insert	· - · - · - · - · - · - · - · - · · - · · - ·
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